

# SEQUENCE LISTING

<110> Yang, Shumin  
 McCall, Catherine A.  
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<120> CANINE AND FELINE IMMUNOREGULATORY PROTEINS, NUCLEIC  
 ACID MOLECULES, AND USES THEREOF

<130> IM-2-C1-C1

<140> not yet assigned

<141> 2001-01-01

<150> 01 312,409

<151> 1999-05-28

<150> 01 087,306

<151> 1993-05-29

<160> 21

<170> PatentIn Ver. 2.1

<210> 1

<211> 16

<212> DNA

<213> Artificial Sequence

<220>

<221> Description of Artificial Sequence: Synthetic  
 Primer

<400> 1

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16

<210> 1

<211> 42

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<213> Artificial Sequence

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<221> Description of Artificial Sequence: Synthetic

<222> Description of Artificial Sequence: Synthetic



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 Gln Asn Leu Ser Leu Ile Lys Glu His Ile Glu Arg Gln Lys Lys Arg  
 90 95 100

tgt gca gga gaa aga tgg aga gtg aca aag ttc cta gag tac ctg caa 388  
 Cys Ala Gly Glu Arg Trp Arg Val Thr Lys Phe Leu Asp Tyr Leu Gln  
 105 110 115 120

gta ttt ctt ggt gta ata aac acc gag tgg aca cgg gaa agt 430  
 Val Phe Leu Gly Val Ile Asn Thr Glu Trp Thr Pro Glu Ser  
 125 130

tgagaacaaa cgggttatt gtagtggaag attttggaga agaattggttt ttggtgatg 490

agaatgaggg ccaaccaaca gtagggactt aatggccagt ataactaagc ttcagagaca 550

aagtaaatat ttcaggcatc ctactacttt atcacttcac acagatgaaa tatatttgag 610

<210> 5

<211> 134

<212> PRT

<213> Canis familiaris

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Met Arg Met Leu Leu Asn Leu Ser Leu Leu Ala Leu Gly Ala Ala Tyr  
 1 5 10 15

Val Ser Ala Phe Ala Val Glu Asn Pro Met Asn Arg Leu Val Ala Glu  
 20 25 30

Thr Leu Thr Leu Leu Ser Thr His Arg Thr Trp Leu Ile Gly Asp Gly  
 35 40 45

Asn Leu Met Ile Pro Thr Pro Glu Asn Lys Asn His Gln Leu Cys Ile  
 50 55 60

Lys Glu Val Phe Gln Gly Ile Asp Thr Leu Lys Asn Gln Thr Ala His  
 65 70 75 80

Gly Glu Ala Val Asp Lys Leu Phe Gln Asn Leu Ser Leu Ile Lys Glu  
 85 90 95

Glu Trp Thr Pro Glu Ser

130

&lt;210&gt; 6

&lt;211&gt; 610

&lt;212&gt; DNA

&lt;213&gt; Canis familiaris

&lt;400&gt; 6

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 catcgcacaa aaaccattct cctccaaaat ctccactac aataagccgg ttggtctca 180  
 actthccggg gtccactcgg tgtttattac accaagaaat acttgcaggt agtctaggaa 240  
 ctthctcact ctccatcttt ctctgcaca ccttttttt tggcgcctta tgtgtcttt 300  
 tattaaagac aagttctgga atagtttata cacagcctcc ccgtgggcag ttggttctt 360  
 caatctgtct atacctgaa aaaattcttt aatgcacagt tggtagtttt tttttcagg 420  
 agtaggaatc atcaggttcc catgcctat cagccaaagt cgatgagtgg agagcagtg 480  
 caagttctct gccaccagtc tttcatggg attttctaca gcaaaggcag aaacataggc 540  
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 gtttcccttg 610

&lt;210&gt; 7

&lt;211&gt; 402

&lt;212&gt; DNA

&lt;213&gt; Canis familiaris

&lt;400&gt; 7

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&lt;210&gt; 8

&lt;211&gt; 400

&lt;212&gt; DNA

&lt;213&gt; Canis familiaris

&lt;400&gt; 8

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ctttgtact ctccatcttt ctctgcaca cctttttttt tggcgctcta tgtgttcttt 120  
 tattaagac aagtttttga atagtttate cacagctcc cgtgggcag ttgggttctt 180  
 caat tgtct ataccctgaa aaacttcttt aatgcacagt tggtgatttt tattttcagg 240  
 agtaagaate atcaggttcc catcgcttat cagccaagtt ccatgagtgg agagcagtgt 300  
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<210> 9

<211> 345

<212> DNA

<213> Canis familiaris

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<221> CDS

<222> (1)..(345)

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 1 5 10 15

ctg ctc tcc act cat cga act tgg ctg ata ggc gat ggg aac ctg atg 96  
 Leu Leu Ser Thr His Arg Thr Trp Leu Ile Gly Asp Gly Asn Leu Met  
 20 25 30

att cct act cct gaa aat aaa aat cac caa ctg tgc att aaa gaa gtt 144  
 Ile Pro Thr Pro Glu Asn Lys Asn His Gln Leu Cys Ile Lys Glu Val  
 35 40 45

ttt cag ggt ata gac aca ttg aag aac caa act gcc cac ggg gag gct 192  
 Phe Gln Gly Ile Asp Thr Leu Lys Asn Gln Thr Ala His Gly Glu Ala  
 50 55 60

ctc cat gaa cta ttc caa aac ttc tct tta ata aac caa cac ata cag 240  
 Val Asn Lys Leu Phe Gln Asn Leu Ser Leu Ile Lys Glu His Ile Glu  
 65 70 75

cgc caa aac aaa agg tgt gca gga gaa aga tgg aga ctg aca aag ttc 288  
 Arg Gln Lys Lys Arg Cys Ala Gly Glu Arg Trp Arg Val Thr Lys Phe  
 85 90 95

ctc cat tct ctg caa cta ttc ctt tgt tta ata aac aac gat tct aca 336  
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115

<210> 10

<211> 115

<212> PRT

<213> Canis familiaris

<400> 10

Phe Ala Val Glu Asn Pro Met Asn Arg Leu Val Ala Glu Thr Leu Thr  
1 5 10 15

Leu Leu Ser Thr His Arg Thr Trp Leu Ile Gly Asp Gly Asn Leu Met  
20 25 30

Ile Pro Thr Pro Glu Asn Lys Asn His Gln Leu Cys Ile Lys Glu Val  
35 40 45

Phe Gln Gly Ile Asp Thr Leu Lys Asn Gln Thr Ala His Gly Glu Ala  
50 55 60

Val Asp Lys Leu Phe Gln Asn Leu Ser Leu Ile Lys Glu His Ile Glu  
65 70 75 80

Arg Gln Lys Lys Arg Cys Ala Gly Glu Arg Trp Arg Val Thr Lys Phe  
85 90 95

Leu Asp Tyr Leu Gln Val Phe Leu Gly Val Ile Asn Thr Glu Trp Thr  
100 105 110

Pro Glu Ser  
115

<210> 11

<211> 148

<212> DNA

<213> Canis familiaris

<400> 11

actttccggt gtcactcgg tgtttattac accaagaagt acttgccggt agtctaggaa 60  
ctttgcact ctccatcttt ctcctgcaca cctttttttt tggcgctcta tgtgttcttt 120

<210> 12

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 12

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36

<210> 13

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 13

cccgccggcgcg ctcaactttc cgggtgtccac tc

32

<210> 14

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 14

accccaaacac tgaacatttc

29

<210> 15

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 15

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<210> 16

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 16

tcaaggagg ctataaattc

20

<210> 17

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 17

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20

<210> 18

<211> 1658

<212> DNA

<213> Canis familiaris

<220>

<221> Intron

<222> (171)..(373)

<220>

<221> Intron

<222> (677)..(1175)



<400> 18

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gattccattat caataatgaa gtaatgagtg ttaataatat ataatgggta accatgctac 300
tcagaagaat tatattaaaa gttatgaacc ttacaataga ttaaaaaatga atgttgtctc 360
ctttcttttt tagaacctga tgattctcac tcttgaaaat aaaaaatgta gttaaattat 420
gatttgataa aatgattaca tgaatcagtt tcatatttta agctataaag tatcagttaa 480
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caagtgggac agjccctttt ttgatgttgt cagttctcca tctcaaaagag ccttgtgtca 660
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agaacaaaac ggttatttgt agtggaaagt ttgggaga 1658
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<210> 19

<211> 1653

<212> DNA

cll-1 clone 1.611.11.11

<400> 19

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ctcttgaca cctttcttgt taaaagaca gaggagatat tgtaggaaac tgtcaattct 180
tgtaaaaaaa aaaaaaaaaa acagggcagag tcagacaaat ataettaagt ttttgcacaa 240
tgtcttaac ttaattttt degatctatg tctctctta ttaaaagaaa gttttgaaat 300
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taatgatata acattttttaa ctttttaata tgaccatgac atactccaga tgtatatgat 780
tccatttagag ttaaaaaatca tttaaataatg tggaagtggac tattaatata ggacttccact 840
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tcagaaaaac gttgaggtga ggaagaccac cgcagaggtt gaatggagcc taggaagtat 960
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attgtaaggt ccataacttt taatataatt cttctgagta acatgggtac ccattatat 1380
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caattacca tgcattttta agaactgtag aatcaaaaag aaaattaccc catcgccat 1500
cagccaagtt cgtatgagtgg agagcagtggt caaggtctct gccaccagtc tattcatggg 1560
attttctaga gcaaaggcag aaacataggt agccccaaga gctagcaaac tcaaattcag 1620
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<210> 20

<211> 15

<212> PPT

<213> Artificial Sequence

<220>

<221> Description of Artificial Sequence: N-terminal  
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5

10

15

<210> 21

<211> 171

<212> RNA

<213> Gene families

<400> 21

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tttgatctcc tgaactcttt aaaaacacac ccttaattcc ttgtgttact attttttaa 240
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